

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U-S PATENT AND TRADEMARK OFFICE WEBSITE SEE BELOW FOR ADDRESS

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1 EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):

 "U.S. Patent and Trademark Office, 220-20th Street-S., Customer Window, Mail Stop Sequence, Crystal-Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER, 091927 071
ATTN: NEW RULES CASES	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS WHICH WERE INSERTED BY PTO SOFTWARE The number/text at the end of each line "present" days and the series of the s
IWrapped Nucleics	The number and a state of the same of the
ahlen Vunuos	was reflieved in a word processor after creating it. Please adjust your right margin to .3; this will
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers:
entropy we have an order of the second of th	he submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please
c:	equence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, sidue having variable length and indicate in the <220>-<223> section that some may be missing
"bug" sc	"bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from anino acid evicusly coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the relevant coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the tificial or Unknown sequences.
Skipped Sequences Sec (OLD RULES) (2) (i) (xi) Thi	Number (s) missing. If intentional, please insert the following lines for each skipped sequence. INFORMATION FOR SEQ ID NO. X. (insert SEQ ID NO where "X" is shown) SEQUENCE CHARACTERISTICS. (Do not insert any subheadings under this heading.) SEQUENCE DESCRIPTION SEQ ID NO. X. (insert SEQ ID NO where "X" is shown)
Plca	se also adjust the "(ii) NUMBER OF SEQUENCES "response to include the skipped sequences
(NEW RULES) <210	pence(s) missing. If intentional, please insert the following lines for each skipped sequences > sequence id number > sequence id number
9 Use of n's or Xaa's Use of	of n's and/or V
(NEW RULES) Per I	of n's and/or Xaa's have been detected in the Sequence Listing 823 of Sequence Rules, use of <220> <223> is MANDATORY if n's or Xaa's are present 20> to <223> section, please explain location of n or Xaa and which are to
10lnyalid <213> Pcr 1.	20> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represent 823 of Sequence Rules, the only valid <213> responses are. Unknown, Artificial Sequence, or ficial Sequence.
Usc 01 < 220>	and the second of the second o
Patentin 2.0 Please resultin listing).	cown." Please explain source of genetic material in <220> to <223> section. Gederal Register, "00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules) do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
n can	only represent a single nucleotide; "Xaa" can only represent a single amino acid
	AMC - Biotechnology Systems Branch - 09/09/2003



IFW16

RAW SEQUENCE LISTING

DATE: 11/15/2004

PATENT APPLICATION: US/09/937,076

TIME: 14:32:50

Input Set : A:\110.01270101.ST25.txt
Output Set: N:\CRF4\11152004\I937076.raw

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3 <110> APPLICANT: MCCARTHY, James B.
         MILESKI, William J.
         JAMIESON, Jr., Gordon A.
         LOW, Walter C.
         SAWCHUCK, Ronald J.
         FURCHT, Leo T.
8
10 <120> TITLE OF INVENTION: METHODS OF USE OF BETA1-INTEGRIN INHIBITORS
12 <130> FILE REFERENCE: 110.01270101
14 <140> CURRENT APPLICATION NUMBER: 09/937,076
15 <141> CURRENT FILING DATE: 2002-03-27
17 <150> PRIOR APPLICATION NUMBER: PCT/US00/07680
18 <151> PRIOR FILING DATE: 2000-03-22
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20 <150> PRIOR APPLICATION NUMBER: 60/125,634
21 <151> PRIOR FILING DATE: 1999-03-22
23 <150> PRIOR APPLICATION NUMBER: 60/167,538
24 <151> PRIOR FILING DATE: 1999-11-24
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Detrualid Response

Pls explain source of genetic

Pls explain source of materiac.

Pls explain source of materiac.
26 <160> NUMBER OF SEQ ID NOS: 18
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33 <213> ORGANISM: ARTIFICIAL
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54 Trp Gln Pro Pro Arg Ala Ala Ile Tyr
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ernor
Summary
55 1
58 <210> SEQ ID NO: 3
59 <211> LENGTH: 8
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61 <213> ORGANISM: ARTIFICIAL
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63 <220> FEATURE:
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error Summary Sheet.

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/937,076

DATE: 11/15/2004 TIME: 14:32:50

Input Set : A:\110.01270101.ST25.txt
Output Set: N:\CRF4\11152004\1937076.raw

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RAW SEQUENCE LISTING DATE: 11/15/2004
PATENT APPLICATION: US/09/937,076 TIME: 14:32:50

Input Set : A:\110.01270101.ST25.txt
Output Set: N:\CRF4\11152004\1937076.raw

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Same errors

RAW SEQUENCE LISTING

DATE: 11/15/2004

PATENT APPLICATION: US/09/937,076

TIME: 14:32:50

Input Set : A:\110.01270101.ST25.txt Output Set: N:\CRF4\11152004\1937076.raw

220 <400> SEQUENCE: 14 222 Ala Arg Ile Tyr 223 1 PIS explain source
of genetic materials
Tryalid
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 11/15/2004 PATENT APPLICATION: US/09/937,076 TIME: 14:32:51

Input Set : A:\110.01270101.ST25.txt
Output Set: N:\CRF4\11152004\1937076.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, oer 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18

VERIFICATION SUMMARY

DATE: 11/15/2004

PATENT APPLICATION: US/09/937,076

TIME: 14:32:51

Input Set : A:\110.01270101.ST25.txt
Output Set: N:\CRF4\11152004\1937076.raw